

っている。RAF キナーゼ、VEGFR-1-3、PDGFR-βなどを標的とするマルチキナーゼ阻害薬 sorafenib のプラセボコントロールによる無作為化比較試験(SHARP trial)の結果、sorafenib 治療群は無治療群に比べ初めて生存期間の延長が確認された。わが国では肝細胞がんに対する第I相試験が行われ、欧米と同じ推奨用量が示されている。今後 sorafenib による化学療法は肝外転移を有する例や局所治療が適応にならない進行例に対する標準治療として位置づけられるものと考えられる。また、肝細胞がんは切除や局所壊死療法など局所治療成功例でも再発率が極めて高いことから、sorafenib は局所治療の補助療法としての適応も期待され、大規模な臨床試験が行われている。さらに現在、より有効な全身治療の確立に向けて、様々な新しい分子標的薬や併用療法の臨床試験が進みつつある。またこれまで sorafenib をはじめ多くの分子標的薬は Child-Pugh A の肝機能が良好な患者でのみ臨床試験が行われてきた。Child-Pugh B 等肝機能低下例での有効性及び安全性を検証する研究も必要と考えられる。

E. 結論

肝外転移を有する肝細胞がんの予後は極めて不良であり、有効な全身化学療法の標準治療の確立が必要である。

F. 健康危険情報

特になし

G. 研究発表

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H. 知的財産権の出願・登録状況

(予定を含む)

特許取得:なし
 実用新案登録:なし
 その他 :なし

Table. Multivariate analysis of prognostic factors in patients with extrahepatic metastasis of hepatocellular carcinoma

Variables		Hazard ratio	95% CI	p-value
Performance status	0			<0.0001
	1-3	2.93	2.07-4.29	
Child-Pugh classification	A			<0.0001
	B/C	2.24	1.62-3.24	
Metastatic organ	Single			0.0022
	Multiple	1.77	1.23-2.55	
No. of metastatic tumor	Solitary			0.037
	Multiple	1.72	1.03-2.86	

研究成果の刊行に関する一覧表

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Original Article

Double-dose double-phase use of second generation hepatitis B virus vaccine in patients after living donor liver transplantation: Not an effective measure in transplant recipients

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Aims: Post-transplant active immunization for chronic hepatitis B patients has been attempted in several studies with controversial results. We assessed the effect of a double-dose double-phase vaccination regimen among partial living donor liver recipients.

Methods: Eighteen patients who underwent liver transplantation (LT) for chronic hepatitis B and two non-hepatitis B virus (HBV)-infected patients who received hepatitis B core antibody (HBcAb)-positive donor organs were recruited 18-78 months after LT. All were on hepatitis B immunoglobulin (HBIG) mono-prophylaxis before and throughout vaccination, to maintain hepatitis B surface antibody (HBsAb) titers of more than 100 IU/mL. Recombinant hepatitis B surface antigen vaccine (40 µg) was administered intramuscularly during weeks 0, 4, 8, 24, 28 and 32.

Results: The patients consisted of 15 males and five females with a median age of 52 (39-59) years. None developed a

sufficient HBsAb titer above 500 IU/mL by week 48. In two patients whose maximum HBsAb titer increased to above 300 IU/mL, we attempted to skip HBIG, but shortly thereafter the titer dropped below 100 IU/mL and HBIG administration was resumed. Although the HBIG dose was reduced during and after vaccination, cessation of administration was not achieved.

Conclusion: Double-dose double-phase use of second generation recombinant vaccine was not effective in this study population. The selected population should be targeted for a conventional vaccine regimen, and different approaches, such as strong adjuvant or pre-S containing protein, should be further tested in a larger number of patients after LT for chronic hepatitis B.

Key words: Hepatitis B vaccine, HBcAb positive donor, HBIG, lamivudine, liver transplantation, prophylaxis

INTRODUCTION

THE LONG-TERM use of hepatitis B immunoglobulin (HBIG) and/or nucleos(t)ide analog prophylaxis has dramatically improved survival rates after liver transplantation for hepatitis B virus (HBV)-related liver disease.¹ Historically, hepatitis B (HB) recurs in approxi-

mately 80% of liver transplant recipients with HBV-related liver diseases. The use of HBIG mono-prophylaxis has improved the rate of recurrent hepatitis to 35%.² Long-term use of HBIG therapy, however, is costly and there is insufficient evidence regarding the optimal length of administration. Lamivudine (LAM) mono-prophylaxis is less costly and is useful for decreasing the rate of hepatitis recurrence to less than 40%.^{3,4} Emerging resistant strains are a concern in long-term follow-up for transplant recipients, however, and additional nucleos(t)ide analogs such as adefovir and entecavir are required.^{5,6} The combination of HBIG and antiviral agents has decreased the rate of hepatitis B recurrence.^{7,8}

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Active immunization against hepatitis surface antigens has been attempted in patients after liver transplantation for HB-related liver diseases.^{9–11} Repeated vaccination effectively initiates the production of anti-hepatitis B antibodies and is thus followed by HBIG administration withdrawal.^{9–11} This idea is theoretically a more economical and simple method compared to passive immunization or nucleos(t)ide analog administration. Contradictory results have been reported, however, and suitable candidates for this type of vaccination have not been determined.^{12–15}

At the University of Tokyo, we use HBIG monophylaxis.¹⁶ The ultimate goal of vaccination is to achieve sufficient production of anti-hepatitis B immunoglobulin and to discontinue further prophylaxis against recurrent hepatitis B. In the present study, we report the results of an active immunization protocol in chronic hepatitis B-related living donor liver transplantation (LDLT) recipients.

METHODS

Subject selection

PATIENTS WHO UNDERWENT LDLT at least 18 months before for HBV-related end-stage liver disease or received hepatitis B core antibody (HBcAb)-positive donor livers were enrolled if they were being treated with an HBIG prophylaxis protocol, free of nucleos(t)ide analogs, without co-infection with hepatitis C virus or human immunodeficiency virus, and if they had no evidence of HBV reactivation. Twenty Japanese patients provided informed consent to the study protocol. There were 15 men and five women with a median age of 52, ranging from 39 to 59 years. Etiologies of end-stage liver disease included chronic hepatitis B in 16, fulminant hepatic failure with a history of chronic hepatitis B in two, and end-stage liver disease due to autoimmune hepatitis and primary biliary cirrhosis in one each. Two non-HBV patients and one chronic hepatitis B patient received core antibody-positive donor organs. Nine patients with hepatocellular carcinoma were free from post-transplant recurrence with a median follow-up period of 43 (18–90) months.

The donors were 13 men and seven women ranging in age from 18 to 54 years and weighing 43–75 kg. Their relationship to the patients included 10 children, five spouses, three nephews, one sibling and one cousin. Right liver graft was performed in eight, extended right graft in one, right lateral graft in two, left lobe with

caudate graft in seven and left lobe graft in two. Three donors were positive for both HB surface antibody (HBsAb) and HBcAb.

Pre- and post-LDLT follow-up protocol

The post-transplantation immunosuppression regimen consisted of steroid induction with tacrolimus, or cyclosporine in case of tacrolimus intolerance, for maintenance.¹⁷ Among the 18 patients with hepatitis B virus infection, LAM 100 mg/day was given orally prior to LDLT. One patient received LAM for more than 1 year, one for 3 months, and 15 received LAM for less than 4 weeks. Of the 17 patients on LAM, a negative HBV-DNA load was confirmed preoperatively in nine cases. In eight patients whose HBV-DNA was detectable pre-transplant, LAM therapy was continued for 4 weeks after LDLT, and discontinued after confirming negative HBV-DNA. Postoperatively, HBIG (Mitsubishi Tanabe Pharma, Hebsulin-IH, Tokyo, Japan) was administered to HBV-infected patients and those who received HBcAb-positive donor organs. Details of the HBIG administration protocol and doses are described elsewhere.¹⁶ In brief, HBIG was administered to maintain the anti-HB surface antibody (HbsAb) levels at greater than 1000 IU/L for patients with HBV and greater than 500 IU/L for patients that received HBcAb-positive donor organs. After 1 year, 1000–2000 U was given intravenously indefinitely to maintain HbsAb levels of greater than 100 IU/L.^{16,18}

Vaccination protocol

After obtaining informed consent, baseline laboratory tests were performed in all patients. Table 1 shows the baseline patient characteristics and laboratory findings. HBV-DNA was undetectable in all patients. Increased dose, namely "double-dose", of recombinant anti-HB vaccination (40 µg/2 mL; Heptavax II, Banyu Pharm, Tokyo, Japan) was injected bilaterally into the deltoid muscles (half dose each side). HBsAb titers were measured at week 0 and every 4 weeks thereafter. Vaccination was scheduled for two phases of three administration cycles at weeks 0, 4, 8, 24, 28 and 32.

During the protocol period, HBIG was administered 2 weeks before and after vaccination if necessary; either 1000 or 2000 IU was administered intravenously according to the previously measured HBsAb titer, as long as it was maintained above 100 IU/mL.

Study end-points and ethical considerations

The primary end-point of this study was the vaccine response. A significant increase, that is >500 IU/mL, in

Table 1 Patient characteristics

Factors	Values ^a
Age	52 (39-59) years
Men, women	15, 5
HBV-related hepatitis	18
HBcAb positive donor	3
Immunosuppression (Tac + CS), (CyA + CS)	15, 5
Months from LDLT (median)	45 (17-90) months
Pre-LDLT LAM	17
Duration of LAM administration before LDLT:	15, 1, 1
<1 month, 1-12 months, >12 months	
HBV-DNA positive at LDLT ^b	8
HBsAg titer at entry	138.5 (92-302) IU/mL
Aspartate aminotransferase	18.5 (7-30) IU/mL
Alanine aminotransferase	16 (5-30) IU/mL
Alkaline phosphatase	197 (103-528) IU/mL
γ -glutamyl transferase	33.5 (11-187) IU/mL
Lactate dehydrogenase	178 (99-315) IU/mL
Total bilirubin	0.9 (0.1-1.5) mg/dl

^aValues are number or median (range). ^bHBV-DNA measured by transcription-mediated amplification methods. Lower detectable limit is 3.7 LEG/mL.

CS, corticosteroid; CyA, cyclosporine; HBsAb, hepatitis B surface antibody; HBcAb, hepatitis B core antibody; HBV, hepatitis B virus; LAM, lamivudine; LDLT, living donor liver transplantation; Tac, tacrolimus.

HBsAb 12 weeks after the last vaccination was considered effective. Secondary end-points were changes in the required HBIG dose.

The study protocol was approved by the institutional review board (No. P2005015-11X) and informed consent was obtained from each patient.

Statistical analysis

The number of units of HBIG administered before and during the study protocol was recorded, and the sum of the number of HBIG units administered at -24 to -1 weeks, 0-23 weeks, 24-47 weeks and 48-71 weeks was compared by the Friedman test.

RESULTS

ALL PATIENTS COMPLETED the full vaccination course. Double-dose vaccination was well-tolerated with the only side effect of local pain not requiring analgesics. During the study period, none of the cases developed active hepatitis or rejection. One patient

developed kidney dysfunction during the study and was unable to complete the HBIG administration protocol (case #5).

Pre-, post- and maximum serum HBsAb titers are shown in Table 2. None of the cases developed a sufficient HBsAb titer level of >500 IU/mL (Fig. 1). None achieved cessation of HBIG administration.

Three cases (#2, #3, #20) developed maximum titers of 313, 408 and 469 IU/mL, and HBIG was thus discontinued in two of them (#2 and #3). The titers then decreased below 100 IU/mL in both cases, however, and HBIG administration was resumed (Fig. 2). Case #20 desired to continue HBIG administration despite the elevation of HBsAb titer to 469 IU/mL. Eventually, HBsAb titer dropped to 214 IU/mL and thus the vaccination was considered ineffective.

HBIG administration in four 24-week periods (-24 to -1, 0 to 23, 24 to 47, 48 to 71) in each case are shown in Table 2. In case 11, HBIG administration between weeks 48 and 71 was stopped due to the appearance of HBsAg and HBV-DNA. HBIG doses as a whole, excluding case 11, decreased over time ($P = 0.006$), as illustrated by the box plot (Fig. 3).

After the final vaccination, all patients were followed for a median of 17 (10-18) months. All patients are alive. In two cases (#11 and #13), serum HBsAg and HBV-DNA were observed at 4 and 8 months after the vaccination protocol, and 60 and 58 months after liver transplantation. In these two cases, the minimum HBsAb level was above 100 IU/mL even after our vaccination protocol. These cases are currently on antiviral therapy and are free from signs of active hepatitis.

DISCUSSION

ACTIVE IMMUNIZATION FOR chronic hepatitis B has been attempted in patients after liver transplantation. In early studies, commercially available recombinant vaccine was used in patients receiving HBIG mono-prophylaxis and was effective in 64-80% of patients.^{9,19} This immune response was sustained for a longer follow-up period of 41 (31-85) months in 14 responders.¹⁰ The use of a potential adjuvant in combination with recombinant vaccine is remarkably effective in 80% of post-transplant recipients, and the HBsAb titer was maintained for a long (8-27 months) follow-up period.¹¹ Studies by other groups, however, demonstrated unfavorable results in patients who were either on LAM prophylaxis or HBIG mono-prophylaxis.¹²⁻¹⁴ Several factors are thought to

Table 2 Characteristics, HBsAb titer and dose requirement of HBIG

Case	Age	Sex	Etiology	Donor HBsAb	Pre-LT HBV-DNA	Pre-LT LAM	Timing of vaccination ¹	IS	Pre- HBsAb ²	Max HBsAb ³	End HBsAb ⁴	HBIG administration				Outcome ⁵ (recurrence ⁶)
												-24-1 weeks ³	48-71 weeks ³	0-23 weeks ³	24-47 weeks ³	
1	52	F	PBC	+	Negative	NA	78	Tac	165	253	253	10	7	10	11	104, alive
2	46	F	AIH	+	Negative	NA	51	CyA	92	313	152	3	5	7	3	75, alive
3	54	M	B-PHF	-	Positive	Yes	48	Tac	122	408	136	12	5	8	5	65, alive
4	39	M	B-PHF	-	Positive	Yes	30	Tac	163	227	221	8	6	6	5	56, alive
5	50	M	BLC, HCC	+	Negative	Yes	90	Tac	175	79	48	6	5	6	7	115, alive
6	57	F	BLC	-	Positive	Yes	57	Tac	127	191	107	7	6	9	7	82, alive
7	47	F	BLC	-	Negative	Yes	56	Tac	140	186	132	7	6	6	6	81, alive
8	50	M	BLC, HCC	-	Negative	Yes	55	Tac	101	153	113	5	6	5	4	77, alive
9	50	M	BLC	-	Positive	Yes	51	Tac	117	114	70	7	6	6	6	75, alive
10	51	M	BLC, IICC	-	Positive	Yes	49	Tac	111	293	138	8	5	6	5	74, alive
11	48	M	BLC, HCC	-	Positive	Yes	48	Tac	281	171	131	11	1	10	4	71, alive, rec 60
12	59	M	BLC	-	Positive	Yes	43	Tac	160	176	162	6	8	6	6	69, alive
13	52	M	BLC, HCC	-	Positive	Yes	43	Tac	209	210	132	8	7	6	6	67, alive, rec 56
14	57	F	BLC, HCC	-	Negative	Yes	41	CyA	109	220	220	12	7	10	8	67, alive
15	56	M	BLC, HCC	-	Negative	None	40	Tac	144	188	93	9	10	12	12	60, alive
16	54	M	BLC	-	Negative	Yes	40	Tac	137	190	190	8	6	5	5	60, alive
17	50	M	BLC	-	Negative	Yes	20	CyA	95	262	140	6	6	6	6	38, alive
18	51	M	BLC, HCC	-	Negative	Yes	19	CyA	135	179	115	11	8	10	9	44, alive
19	52	M	BLC, HCC	-	Negative	Yes	18	Tac	152	245	121	14	7	12	6	43, alive
20	53	M	BLC	-	Negative	Yes	17	Tac	302	469	214	14	10	12	12	37, alive

¹Number indicates months after liver transplantation. ²IU/mL. ³Number indicates vials of HBIG (1V = 1000 Units) used during the period. ⁴Number indicates months after liver transplantation, when HBV-DNA became positive.

AIH, autoimmune hepatitis; B-PHF, hepatitis B related fulminant hepatic failure; BLC, hepatitis B related liver cirrhosis; CyA, cyclosporine A; PHF, fulminant hepatic failure; HBsAb, hepatitis B surface antibody; HBcAb, hepatitis B core antibody; HBIG, hepatitis B immunoglobulin; IICC, hepatocellular carcinoma; IS, immunosuppression; LAM, lamivudine; LC, liver cirrhosis; LT, liver transplantation; PBC, primary biliary cirrhosis; Tac, tacrolimus.

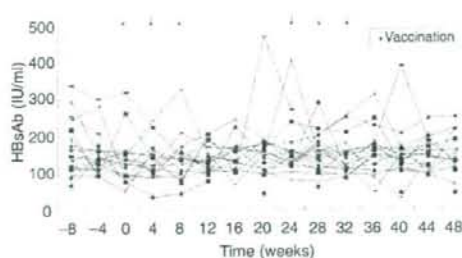


Figure 1 Change in serum hepatitis B surface antibody (HBsAb) titers before and during the vaccination protocol. Heptavax II (40 µg) was administered at weeks 0, 4, 8, 24, 28 and 32. HBsAb levels were between 100 IU/mL and 200 IU/mL in most cases. There was no significant response to vaccination throughout the study.

contribute to a good response, such as younger age, use of an adjuvant to the vaccine and negative HBV-DNA preoperatively. Use of LAM was once speculated to have a negative effect, and, in another study, Angelico *et al.*

failed to reveal a favorable effect of mono-prophylaxis with HBIG.¹²

In our study, the double-dose double-phase use of a second generation recombinant hepatitis B vaccine was tested. Our institution applies an HBIG mono-prophylaxis protocol against hepatitis B recurrence. For non-replicate HBV disease, LAM is discontinued at the time of transplantation. During this study, HBIG was administered throughout the vaccination protocol according to Binzele's report,¹¹ and anti-hepatitis B antibody levels of greater than 500 IU/mL were determined to be effective. Unfortunately, none of the 20 patients developed an adequate anti-hepatitis B antibody titer after two vaccination cycles. Although the dose of HBIG administration decreased during and after the vaccination protocol, we did not consider it a significant change, since none achieved cessation of HBIG administration. The possible factors contributing to vaccination failure among the subjects are that 15 of the 20 patients were aged 50 years or older, eight of 18 chronic hepatitis B patients had a HBV replicative state at the

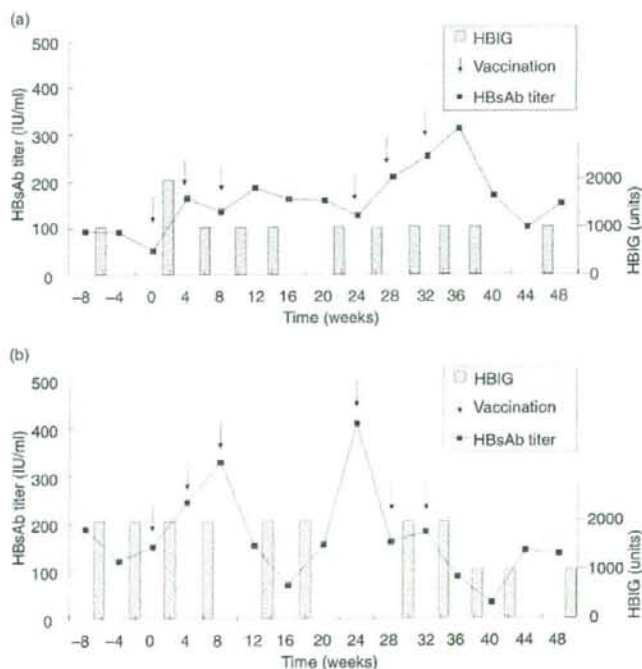


Figure 2 Hepatitis B surface antibody (HBsAb) titers in response to vaccination. (a) Case 2: 46-year-old female with auto-immune hepatitis (AIH), hepatitis B core antibody (HBcAb)-positive donor. (b) Case 3: 54-year-old male with B-fulminant hepatic failure. Hepatitis B immunoglobulin (HBIG) administration was skipped in response to elevated HBsAb level. The HBsAb titer then dropped and HBIG administration was resumed in both cases.

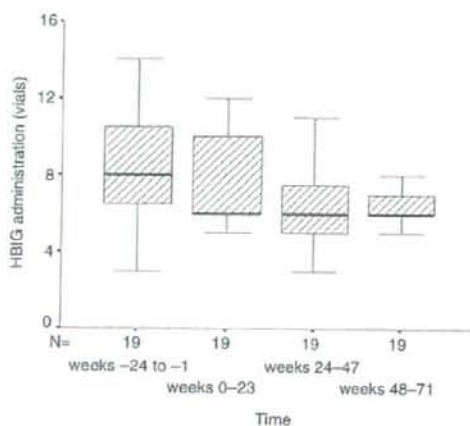


Figure 3 Hepatitis B immunoglobulin (HBIG) administration in four 24-week periods (-24 to -1, 0-23, 24-47, 48-71) is graphically shown by box plot. The dose of HBIG administration decreased over the time period ($P=0.006$). Box plot explanation: upper horizontal line of box, 75th percentile; lower horizontal line, 25th percentile; horizontal bar within box, median; upper horizontal bar outside box, 90th percentile; lower bar outside box, 10th percentile.

time of transplantation, use of corticosteroids in combination with a calcineurine inhibitor, and one patient was receiving hemodialysis.²⁰ These factors might have affected the results of this study negatively.

A second generation recombinant HB vaccine is used for prophylaxis in the healthy population. HBsAg-specific T and B cells are induced and a sufficient amount of HBsAb can be produced to neutralize circulating HB virus particles. The recombinant HB vaccines containing S, pre-S2 and/or pre-S1, the so-called the third generation vaccines, have immunogenic advantages over the second generation recombinant HB vaccines; they more efficiently induce an immune response than the second generation HB vaccines among a healthy population,^{21,22} and induce not only anti-S antibodies, but also anti-pre-S2 antibodies. Such a vaccine containing S, pre-S1 and pre-S2 antigens was used among Chinese patients who underwent liver transplantation for chronic HB and were on LAM prophylaxis.²³ The authors of that study reported that the vaccine was effective in 10 of 20 patients. An earlier study by Karasu *et al.*,¹³ however, failed to show the effectiveness of pre-S1, pre-S2 and S gene products.

Prevention of hepatitis B infection is equally important in HBsAg-negative patients receiving HBe-positive donor organs, as the risk for de novo HB is high.²⁴ The current standard of care for transplant recipients of HBeAb-positive donor organs is similar to that for patients with chronic HB, including long-term HBIG and/or nucleos(t)ide analog.^{16,24-26} In our study, HB vaccine was administered to two HBV non-infected patients who received HBeAb-positive donor organs. Neither of these two patients, however, responded. Pediatric transplantation recipients who receive prior vaccination under an immunization program are likely to achieve a high anti-HB titer by active immunization after LDLT.²⁷ LAM prophylaxis was withdrawn after 2 years if an adequate anti-hepatitis B titer was achieved. Soejima *et al.*²⁸ recently reported that 6 of 11 Japanese patients receiving liver transplants for fulminant hepatitis B or for non-HBV diseases with HBeAb-positive donor organs had seroconversion. The tested patients were younger, with a median age of 33 years. These patients were on combination prophylaxis with LAM and HBIG, and HBIG was withdrawn after vaccination. The question remains, however, as to whether LAM therapy can be discontinued in the future among this population.

Vaccination may be promising in selected populations, such as in younger recipients or in those with fulminant hepatitis or HBsAg-negative recipients receiving HBeAb-positive donor organs. For patients older than 50 with vaccination failure or chronic hepatitis B patients, a different approach may prove optimal, such as the use of a pre-S containing vaccination. The research findings for the use of such vaccines are controversial, however, warranting further study.

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Long-term trends of the incidence of hepatocellular carcinoma in the Nagasaki prefecture, Japan

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Abstract. The incidence of hepatocellular carcinoma (HCC) in Japan is still increasing. The aim of the present study was to analyze the epidemiological trend of HCC in the Western area of Japan, Nagasaki. A total of 1,807 patients with HCC diagnosed at our two hospitals between 1981 and 2005 were consecutively recruited for this study. Cohorts of patients with HCC were categorized into five-year intervals. The etiology of HCC was categorized into four groups: HCC-B: HBsAg positive, HCVAb negative, HCC-C: HCVAb positive, HBsAg negative, HCC-BC: both of HBsAg and HCVAb positive and HCC-nonBC: both of HBsAg and HCVAb negative. The number and proportion of HCC-B cases decreased from 1986 to 1990 and thereafter stabilized, whereas those of HCC-C reached the peak from 1995 to 2000 and thereafter decreased. On the other hand, the number and ratio of the HCC-nonBC cases continued to increase in the whole period. The male/female ratio of HCC-C patients decreased from 6.4 in the period 1981-1985 to 1.9 in 2001-2005, indicating clearly the increase of female patients. On the other hand, the male/female ratio of other types of HCC patients did not change during the period. HCC patients rapidly increased from 1981 to 2000 and this increase was originated from that of HCC-C. The increase of the median age and the number of female patients with HCC-C was also demonstrated. The increase in the number and the proportion of the HCC-nonBC patients was also significant.

Introduction

Primary liver cancer is the most common primary cancer of the liver accounting for ~6% of all human cancers. It is estimated that half a million cases occur worldwide annually, making

primary liver cancer the fifth most common malignancy in men and the ninth in women (1-6). Hepatocellular carcinoma (HCC) accounts for 85 to 90% of primary liver cancers (7) and the age-adjusted HCC mortality rate has increased in recent decades in Japan (8). Similarly, a trend of increasing rates of HCC has been reported from several developed countries in North America, Europe and Asia (9,10). HCC often develops in patients with liver cirrhosis caused by hepatitis B virus (HBV), hepatitis C virus (HCV), excessive alcohol consumption, or nonalcoholic fatty liver disease. Of the hepatitis viruses which cause HCC, HCV is predominant in Japan (11-14).

Although the age-adjusted incidence of HCC has increased in Japan, sequential changes in background features of HCC patients are not fully understood (15). Yoshizawa reports that deaths due to HCC in Japan have continued to increase in males, particularly in those older than 60 years of age in the past 3 decades, although the reasons for this are unclear (16). To clarify factors affecting epidemiological changes in Japanese HCC patients, especially the change in age distribution and gender, we analyzed the underlying features of HCC patients in a two major liver center-based study.

Patients and methods

Patients. A total of 1,807 patients with HCC diagnosed between January 1981 and December 2005 in the Liver Disease Center, National Nagasaki Medical Center and in the outpatient clinic of The First Department of Internal Medicine, Nagasaki University Hospital, were consecutively recruited for this study. The diagnosis of HCC was based on AFP levels and imaging techniques including ultrasonography (USG), computerized tomography (CT), magnetic resonance imaging (MRI), hepatic angiography (HAG) and/or tumor biopsy. The diagnostic criteria for HCC were either a confirmative tumor biopsy or elevated AFP (≥ 20 ng/ml) and neovascularization in HAG and/or CT. Cohorts of patients with HCC were categorized into five-year intervals (1981-1985, 1986-1990, 1991-1995, 1996-2000 and 2001-2005).

Etiology of HCC. Sera were stored at -80°C until use. A diagnosis of chronic HCV infection was based on the presence of HCVAb (microparticle enzyme immunoassay; Abbott

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Key words: hepatitis C virus, hepatocellular carcinoma, aging, Japan

Table I. The characteristics of HCC patients, 1981-2005.

Period	1981-1985	1986-1990	1991-1995	1996-2000	2001-2005	Total
Number	240	316	369	419	463	1807
Gender						
Male	194	257	268	314	314	1347
Female	46	59	101	105	149	460
Ratio (male/female)	4.2	4.4	2.7	3.0	2.1	2.9
Age (y.o) (IQR)	57 (6.5)	61 (5.1)	63 (5.4)	66 (5.1)	68 (6.3)	64 (6.5)
Hepatitis virus						
HCC-B	95	70	80	67	100	412
HCC-C	111	213	240	292	278	1134
HCC-B+C	8	8	9	11	10	46
HCC-nonBC	26	25	40	49	75	215

Gender: 2000-2005 vs. 1981-1985 $p=0.0003$; 2000-2005 vs. 1986-1990 $p\leq 0.0001$; 2000-2005 vs. 1991-1995 $p=0.1330$; 2000-2005 vs. 1996-2000 $p=0.0197$. Age: 2000-2005 vs. 1981-1985 $p\leq 0.0001$; 2000-2005 vs. 1986-1990 $p\leq 0.0001$; 2000-2005 vs. 1991-1995 $p\leq 0.0001$ and 2000-2005 vs. 1996-2000 $p=0.0292$. IQR, interquartile range.

Laboratories) and HCV-RNA detected by polymerase chain reaction (PCR), whereas diagnosis of chronic HBV infection was based on the presence of hepatitis B surface antigen (HBsAg) (enzyme-linked immunosorbent assay; Abbott Laboratories).

Statistical analysis. The data were analyzed by the Mann-Whitney test for the continuous ordinal data between two qualitative variables. The standard deviation was calculated based on the binomial model for the response proportion. $P<0.05$ was considered statistically significant.

Results

Clinical features of the studied patients. A total of 1,807 patients with HCC were diagnosed at our hospital from 1981 to 2005. There were 1,347 male (75%) and 460 female (25%) patients, with a median age of 64 years. The proportion of patients diagnosed as HCC-B (HBV-associated: HBsAg positive, HCVAb negative) was 23% (412 of 1,807), whereas 63% (1,134 of 1,807) had HCC-C (HCV-associated: HCVAb positive, HBsAg negative) and an additional 3% (46 of 1,807) had HCC associated with both viruses. The remaining 215 patients (12%) showed both of the virus markers negative.

As shown in Table I and Fig. 1, the number and proportion of HCC-B cases decreased from 1986 to 1990 and thereafter stabilized, whereas those of HCC-C reached the peak in the period 1996-2000 and thereafter decreased. The number and proportion of the HCC-nonBC (HBsAg and HCVAb negative) cases continued to increase in the whole period.

Background features for patients with HCC. Fig. 2 shows the median age at diagnosis of HCC-B, HCC-C and HCC-nonBC in five-year intervals (1981-1985, 1986-1990, 1991-1995, 1996-2000 and 2001-2005). The median age of patients at diagnosis of HCC-C showed a steadily significant increase

from 58 to 69 years of age during the period. The median age of patients with HCC-B and HCC-nonBC did not significantly change during the period.

Fig. 3 shows the age distribution of patients with HCC-B and HCC-C with the five 5-year intervals. There was no difference in the age distribution of patients with HCC-B during these periods. In contrast, HCC-C obviously had a trend to increase in the number of patients aged >65 years.

Table I shows that the male/female ratio of HCC patients decreased from 4.2 in the period 1981-1985 to 2.1 in 2001-2005, indicating clearly the increase of female patients. In analysis of patients in HCC-C, the male/female ratio in the periods 1981-1985, 1986-1990, 1991-1995, 1996-2000 and 2001-2005 were 6.4, 4.8, 2.5, 2.7 and 1.9, respectively (1981-1985 vs. 2001-2005, $p\leq 0.0001$) (Table II). The ratio became clearly smaller, indicates an increase in female patients with HCC-C. On the other hand, the male/female ratio of other types of HCC patients did not significantly change during the period.

Discussion

This was a two major liver center-based study designed to examine the sequential change in the background of HCC patients during the past 25 years, 1981-2005. More than 80% of our patients had chronic HBV or HCV infections. During the observation period, the number and proportion of HCC-B cases decreased in the period 1986-1990 and thereafter reached a plateau, whereas HCC-C reached a peak in the period 1996-2000 and thereafter slightly decreased. On the other hand, the number and the proportion of HCC-nonBC gradually increased in the periods of 1981-1985, 1986-1990, 1991-1995, 1996-2000 and 2001-2005 being 26 (11%), 25 (8%), 40 (11%), 49 (12%) and 75 (16%), respectively. Previous studies from Japan reported that the proportion of HCC-C had been increased and reached a plateau in the

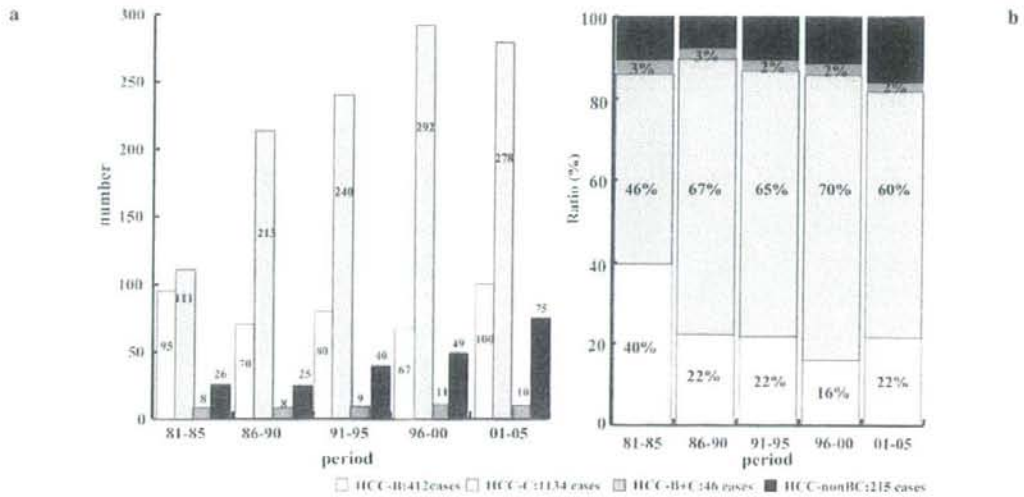


Figure 1. Sequential changes in the number (a) and ratio (b) of HCC patients categorized by etiology during the period 1981-2005 with 5-year intervals.

Table II. The number and male/female ratio of HCC patients during the period of 1981-2005.

Period	1981-1985	1986-1990	1991-1995	1996-2000	2001-2005	Total
Number	240	316	369	419	463	1807
Total						
Male	194	257	268	314	314	1347
Female	46	59	101	105	149	460
Ratio (male/female)	4.2	4.4	2.7	3.0	2.1	2.9
HCC-B						
Male	69	54	61	55	74	313
Female	26	16	19	12	26	99
Ratio (male/female)	2.7	3.4	3.2	4.6	2.9	3.2
HCC-C						
Male	96	176	172	212	182	838
Female	15	37	68	80	96	296
Ratio (male/female)	6.4	4.8	2.5	2.7	1.9	2.8
HCC-nonBC						
Male	21	20	29	40	51	1347
Female	5	5	11	9	24	460
Ratio (male/female)	4.2	4.0	2.6	4.4	2.1	2.9

HBV and nBnC: NS, HCV: 2000-2005 vs. 1981-1985 $p \leq 0.0001$; 2000-2005 vs. 1986-1990 $p \leq 0.0001$; 1996-2000 vs. 1981-1985 $p = 0.0033$; 1996-2000 vs. 1986-1990 $p = 0.0084$; 1991-1995 vs. 1981-1985 $p = 0.0024$ and 1991-1995 vs. 1986-1990 $p = 0.0058$.

period of 1981-2001 (8,15,17-19). However, in our study, the number and proportion of HCC-C cases decreased in the period 2001-2005. This may be due to interferon therapy associated with a decreased incidence of HCC (20-24). Iron depletion for chronic hepatitis C patients is a promising modality for lowering the risk of progression to HCC

(25,26). Oral supplementation with oral branched-chain amino acids has been useful in the prevention HCC (27). Finally, the chronically HCV-infected population is aging in Japan. Yoshizawa reported that age-specific prevalence for the presence of HCVAb among ~300,000 voluntary blood donors from Hiroshima in 1999 clearly increased with the

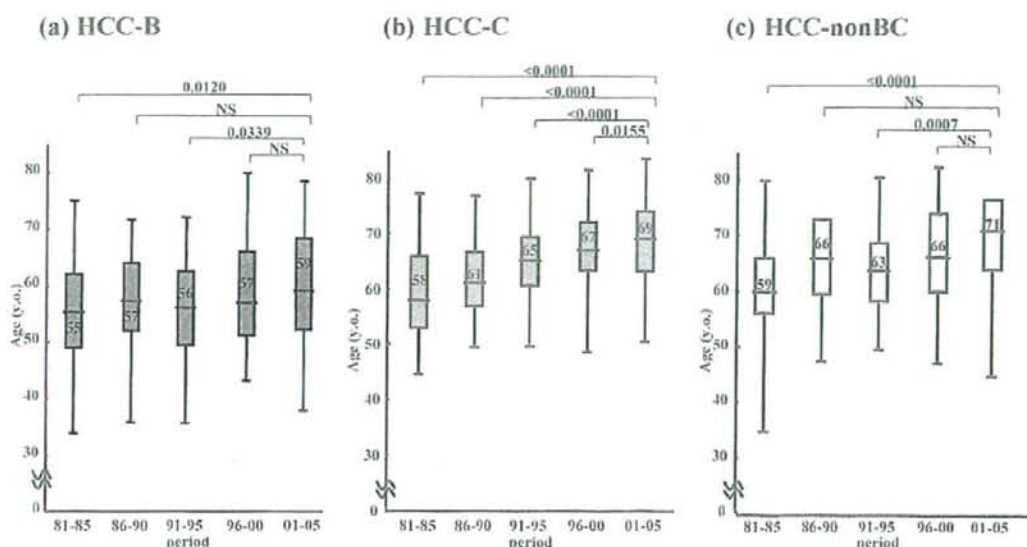


Figure 2. Sequential changes in the median age of HCC patients categorized by etiology during the period, 1981-2005 with 5-year intervals. (a) HCC-B, (b) HCC-C and (c) HCC-nonBC type $p < 0.05$.

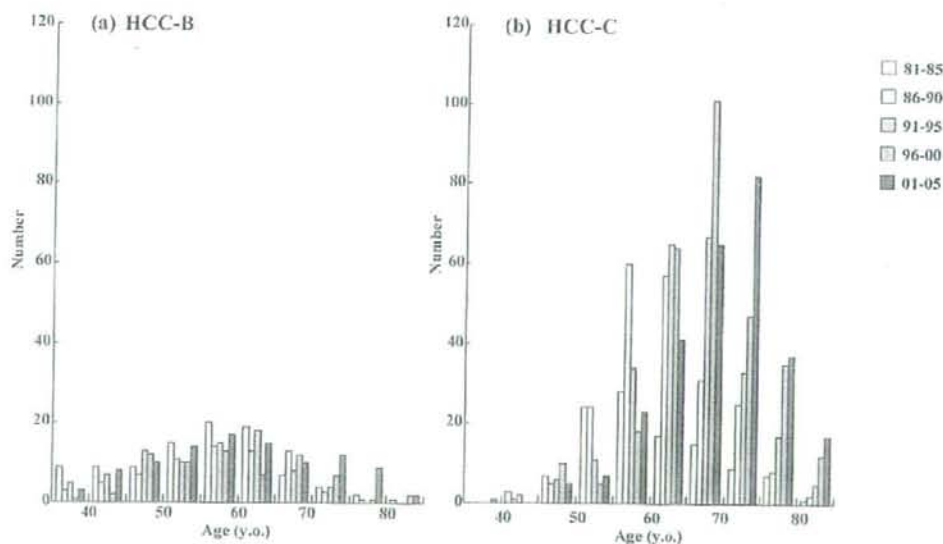


Figure 3. Changes in the age distribution of patients with HCC-B and HCC-C during the period, 1981-2005 with 5-year intervals.

age, reaching the highest proportion of 7% in individuals who were >70 years old (15,16). In this study, the median age of patients with HCC-C steadily increased from 58 to 69 years of age during the studied period. *i.e.* HCV infected people become older and they were regarded as a high risk for HCC.

In almost all populations, males have higher liver cancer proportions than females, with the male/female ratios usually

averaging between 2:1 and 4:1 (7). However, the male/female ratio of HCC in Japan was 4.5 in the period 1983-1985 and 2.57 in 2000-2001 (17). In analysis of background features among HCC patients, HCC-B and HCC-nonBC cases revealed no significant change, whereas the male/female ratio of patients with HCC-C steadily decreased from 6.4 to 1.9 during the period. We suggest that the increase of female

patient with HCC-C was caused by the aging of HCV infected people. The increase of females among HCC patients was considered to increase because of HCC-C.

It is known that 2 to 4 decades of chronic HCV infection are required to develop cirrhosis and subsequent HCC (28-31). The number of HCC cases has increased in Japan, because individuals infected with HCV in the past have grown old and have reached the cancer-bearing age. The prevalence of HCV infection in young Japanese individuals is low and the incidence of HCVAb is very low because of preventative actions against HCV infection such as the screening of blood products for HCV and the use of sterile medical equipment (32). Additionally, we showed that the number and proportion of patients with HCC-C cases decreased together with an increase in the median age, whereas the number and ratio of HCC-nonBC steadily increased during the studied period. Based on these findings it may be expected that the incidence of HCC-nonBC in Japan may continue to increase even after the consequence of the HCV epidemic level off in the near future, although Japan is far advanced with regard to HCC-C.

In summary, HCC patients rapidly increased from 1981 to 2000 and this increase originated from HCC-C and the increase of the median age and the number of female patients with HCC-C. Increase in the number and proportion of the HCC-nonBC patients are also significant.

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Interleukin-18 promoter polymorphisms and the disease progression of Hepatitis B virus-related liver disease

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In this study, we aimed to explore whether interleukin-18 (IL-18) gene-promoter polymorphisms are associated with the outcome of hepatitis B virus (HBV) infection. In all, 204 chronically HBV-infected patients were recruited in this study. Of the 204 HBV-infected patients, 43 were considered to be inactive HBV carriers based on the sustained normalization of serum alanine aminotransferase (ALT) together with seropositivity for the antibody to hepatitis B e-antigen (anti-HBe). A total of 161 patients were found to have chronic progressive liver disease, which included cirrhosis. In these HBV-infected patients, the frequencies of AA genotype of IL-18 gene-promoter polymorphisms at position -607 and C allele at position -137 were significantly higher in inactive HBV carriers compared with those in patients with chronic progressive liver disease. These polymorphisms of the IL-18 promoter regions (-607 and -137) could be associated with different outcomes of HBV infection. (*Translational Research* 2009;153:91-96)

Abbreviation: ALT = alanine aminotransferase; CI = confidence interval; CPLD = chronic progressive liver disease; anti-HBe = hepatitis B e-antigen; HBsAg = hepatitis B surface antigen; HBV = hepatitis B virus; HCC = hepatocellular carcinoma; IFN- γ = interferon- γ ; IL-18 = interleukin-18; OR = odds ratio; PCR = polymerase chain reaction; SNP = single nucleotide polymorphism; Th₁ = T-helper type 1; TMA = transcription-mediated amplification

Hepatitis B virus (HBV) infection is one of the most prevalent chronic viral diseases in the world.¹ Most individuals with persistent HBV in-

fection develop chronic hepatitis, which can progress to cirrhosis or hepatocellular carcinoma (HCC).² Increasing evidence indicates that genetic factors influence the natural history of chronic liver diseases.³ Furthermore, several recent advances concerning the polymorphism of cytokines that control the host response to the virus could play an important role in determining the outcome of HBV infection.⁴ Previous studies have shown that the capacity of cytokine production varies among individuals and correlates with single nucleotide polymorphisms (SNPs) in the promoter region of various cytokine genes.⁵ Also, cytokine gene polymorphisms were associated with the severity of the liver disease in patient with HBV infection,⁶ which may provide clues to clarify the mechanism for the progression of viral hepatitis.

Interleukin-18 (IL-18) was first described as an interferon- γ (IFN- γ)-producing factor, and it has multiple functions including the activation of cytotoxic T lymphocytes or natural killer cells and the promotion

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AT A GLANCE COMMENTARY

Background

The mechanisms underlying pathogenesis and the progression of chronic hepatitis B virus (HBV) infection have not been properly elucidated. Accordingly, no satisfactory therapeutic approaches are available to treat the millions of chronic HBV carriers. Although chronic HBV infection is induced by HBV, the immune system plays an important, yet poorly understood, role in its pathogenesis.

Translational Significance

In this study, we investigated a relationship between interleukin (IL)-18 promoter polymorphisms and the disease progression of chronic HBV-related liver diseases. Our data suggest that polymorphisms at the IL-18 gene-promoter region could be associated with different outcomes of HBV infection.

of T-helper type 1 (Th₁)-type immune responses.⁷ Considering these multiple functions, IL-18 may activate effector cells that are involved in the cytotoxicity against pathogens or malignant cells. Because it is involved in the inflammatory cytokine network, IL-18 could have an important role in the development of chronic inflammatory diseases.⁸ Recent findings show that the IL-18 gene-promoter region regulates the gene expression of this cytokine.⁹ Interestingly, 2 SNPs at position -607 A/C and -137 G/C within the IL-18 promoter region were suggested to alter the IL-18 promoter activity.⁹ Taking this evidence into consideration, we investigated the possible role of the SNPs of IL-18 gene-promoter region in the progression of chronic hepatitis B.

PATIENTS AND METHODS

Patients. A total of 204 patients, who were positive for the hepatitis B surface antigen (HBsAg) and who visited the clinics for treatment of liver disease at Nagasaki University Hospital or at National Nagasaki Medical Center between August 1999, and December 2003, were enrolled in this study. Sixty three healthy volunteers (33 males and 30 females) served as a control group. The patients were followed regularly with measurements of serum alanine aminotransferase (ALT) and HBV markers, such as HBsAg, hepatitis B e-antigen (HBeAg), and anti-HBe, using commercially available radioimmunoassay kits (Dainabot, Tokyo, Japan) every 3–6 months. Patients were also examined with ultrasonography or computed

Table 1. Clinical characteristics of 204 HBV carrier

Variable	Patients with inactive HBV carrier (n = 43)	Patients with CPLD (n = 161)
Sex (male/female)	20/23	114/47
Age (years)	55 ± 18	50 ± 14
HBeAg/anti-HBe status	0/43	53/109
Chronic hepatitis/cirrhosis		68/93
Serum ALT (IU/L)	21 ± 8	74 ± 101
HBV-DNA	(n = 31)	(n = 148)
< 10 ⁵ copies/mL	31	60
≥ 10 ⁵ copies/mL	0	88
Liver biopsy specimen available during study period	(n = 6)	(n = 51)
Stage of fibrosis		
F0–F2	6	15
F3–F4	0	36

NOTES: Age and serum ALT are expressed as mean ± SD.

tomography of the liver every 3–6 months. Serum HBV-DNA was detected by the transcription-mediated amplification (TMA) method¹⁰ as described previously. The results were expressed as the logarithm of the genome equivalent per milliliter (LGE/mL). The detection limit of this method is 3.7 LGE/mL. The patients receiving IFN therapy or antinucleoside analog reverse transcriptase inhibitor therapy were excluded in this study. The study protocol was approved by the Ethics Committees of both the Nagasaki University Hospital and the National Nagasaki Medical Center, and informed consent was obtained from each individual. Of 204 patients, 43 (20 male, 23 female; mean age ± standard deviation [SD], 55 ± 18 years) were considered to be inactive HBV carriers based on sustained normal serum ALT levels together with seropositivity for anti-HBe throughout the study. Of 43 inactive HBV carriers, all 31 patients tested for HBV-DNA had serum HBV-DNA levels less than 10⁵ copies/mL (Table 1). In addition, 161 of the 204 HBV carriers were considered to have chronic progressive liver disease (CPLD) such as chronic hepatitis or cirrhosis, which is manifested by elevated serum ALT levels and by clinical or histologic findings on liver tissue study during the follow-up period. This group comprised 114 males and 47 females (mean age ± SD, 50 ± 14 years). The ratio of males to females was significantly higher in CPLD patients compared with those in inactive HBV carrier ($P = 0.005$) (Table 1). In accordance with the previous report, this ratio demonstrated that the patient gender affects the disease progression in HBV infection.¹¹ The group of patients with CPLD was classified into the following 3 subgroups: (1) 62

Table II. Primers for the SNP analysis

PCR	SNP	Primer	Sequence
1st	-607 (A/C)+ -137 (G/C)	-607ControlF	5'-CTTTGCTATCATTCCAGGAA-3'
		-137R	5'-AGGAGGGCAAATGCCTGG-3'
2nd	-607	-607BtF	5'-biotin-CTTTGCTATCATTCCADGGAAATAGAAAGTTT-3'
		-607BseR	5'-TGCTGTATCAGATGCAAGCCAGACGGATACCATGAGGAGAAATTTAT-3'
2nd	-137	-137BtR	5'-biotin-ACTGCTGTCCGCACTCCTTGGGCCCGC-3'
		-137BseF	5'-GAGGTACAGGTTTTTGAAGGCACAGGCCCACTGAGGAGGAAGAA-3'

patients with chronic hepatitis (CH), (2) 52 patients with cirrhosis, and (3) 47 patients with HCC. Of 47 HCC patients, 41 patients (87%) had cirrhosis.

Of 204 HBV carriers, 57 had undergone liver biopsy during the study period, and their degree of liver fibrosis were assessed using the METAVIR system.¹² To ensure a sufficient number of patients in each category, the severity of fibrosis was classified into 2 categories: F0–F2 and F3–F4. Among 57 patients, 36 patients in the CPLD group were classified as category F3–F4. (Table I).

DNA extraction. Genomic DNA was isolated from whole blood using the QIAamp DNA blood protocol (Qiagen Ltd., West Sussex, United Kingdom) according to the manufacturer's instructions.

IL-18 (-607/-137) genotyping. Two SNPs at position -607 A/C (rs1946518) and -137 C/G (rs187238) in the IL-18 gene promoter were determined by the ligation-mediated genotyping method¹³ with a slight modification. A 849-bp fragment that contained the 2 SNPs was amplified by polymerase chain reaction (PCR) using primers -607ControlF and -137R (Table II), LA Taq polymerase (Takara Bio Inc., Shiga, Japan), and approximately 20 ng genomic DNA. The products were subjected to the 2nd PCR to amplify 178-bp and 174-bp fragments for genotyping at -607 and -137 by the primer pairs of -607BtF and -607BseR and of -137BtR and -137BseF, respectively. The second PCR products for the 2 SNPs were mixed and digested by BseRI to generate a 2-base overhang at the SNP sites. The 2 adapters complementary to the 2 possible 2-base overhang generated from a single SNP were prepared by annealing oligonucleotides, one of which was labeled with 2 different fluorescent dyes (FITC/TexasRed for -607/A or Cy3/Cy5 for -137/G/C) and mixed. The BseRI digest was ligated with the mixed adapters by Ligation Convenience Kit (Nippon Gene Co., Ltd., Toyama, Japan) at room temperature (Table III). The biotinylated DNA fragments were bound to Dynabeads M-280 Streptavidin (DynaL Biotech [Invitrogen], Tokyo, Japan) were subjected to the fluorescence measurement after extensive washing. The reactions after the 2nd PCR were automated using MagSNiPer FD (PSS Co., Ltd., Chiba, Japan) equipped with a 12-channel paramagnetic beads handling unit.^{14,15}

Table III. Oligonucleotides for adapter preparation

SNP (genotype)	Sequence*
-607 (C)	5'-FITC-TACAAGATTCTGAAGACACCACCCAT CCTTGT-3'
-607 (A)	5'-TexasRed-TACAAGATTCTGAAGACACCACCCA TCCTTIT-3'
-137(C)	5'-Cy3-TACAAGATTCTGAAGACACCACCCATCCT TGA-3'
-137(G)	5'-Cy5-TACAAGATTCTGAAGACACCACCCATCCT TGA-3'
Common†	5'-AAGGATGGGTGGTCTTCAGAACTCTGTGA-3'

*The oligonucleotide was annealed to each oligonucleotide above.

†The nucleotides at the position of the SNPs are underlined.

Statistical analysis. Results are expressed as mean \pm SD. Comparison of the allele and genotype frequencies of different groups were performed using the chi-square test or the Fisher exact test. IL-18 allele frequencies were tested for the Hardy-Weinberg equilibrium for both patients and controls using the Cochran-Armitage test. The significance level was set at a *P* value of 0.05.

RESULTS

We investigated the distribution of IL-18 promoter -137 (C/C) and -607 (A/C) polymorphisms in 204 Japanese HBV-infected patients (case) and 63 healthy volunteers (control). The genotypes at the positions of IL-18 promoter -137 (C/C) and -607 (A/C) polymorphisms were in Hardy-Weinberg equilibrium in both the case subjects and the control subjects (Table IV and Table V).

The genotype frequencies of IL-18 promoter polymorphisms (-607 and -137) in each subgroup of HBV-infected patients are summarized in Table VI. These 204 HBV-infected patients were divided into the 2 groups; 43 patients were considered to be the inactive HBV carrier, and 161 were found to have chronic progressive liver diseases (CPLD). The genotype frequencies of IL-18 promoter polymorphisms (-607 and -137) in these 2 groups of HBV carriers are summarized in Table VII. With regard to the -607 genotypes, 6 (14.0%) inactive HBV carriers had the CC genotypes,

Table IV. Frequencies of IL-18 gene-promoter genotypes (-607 and -137) in case subjects

Locus	Genotype	Observed number (%)	Expected number*	P value†
IL-18 -607	A/A	55 (27.0)	64.2	0.46
	A/C	119 (58.3)	105.5	
	C/C	30 (14.7)	39.3	
IL-18 -137	G/G	167 (81.9)	164.1	0.37
	G/C	32 (15.7)	37.9	
	C/C	5 (2.5)	2.2	

*Expected phenotype frequencies based on observed allele frequencies and assuming Hardy-Weinberg equilibrium.

†P values were calculated using the Cochran-Armitage test for Hardy-Weinberg equilibrium at individual loci.

Table V. Frequencies of IL-18 gene promoter genotypes (-607 and -137) in control subjects

Locus	Genotype	Observed number (%)	Expected number*	P value†
IL-18 -607	A/A	20 (31.7)	19.48	0.50
	A/C	30 (47.6)	31.10	
	C/C	13 (20.6)	12.42	
IL-18 -137	G/G	52 (82.5)	51.57	0.48
	G/C	10 (15.9)	10.68	
	C/C	1 (1.6)	0.57	

*Expected phenotype frequencies based on observed allele frequencies and assuming Hardy-Weinberg equilibrium.

†P values were calculated using the Cochran-Armitage test for Hardy-Weinberg equilibrium at individual loci.

19 (44.2%) had the AC genotype, and 18 (41.9%) had the AA genotype. Of the CPLD group, 24 patients (14.9%) had the CC genotype, 100 (62.1%) had the AC genotype, and 37 (23.0%) had the AA genotype (Table VII). The frequency of the AA genotype was significantly lower in CPLD compared with that in the inactive HBV carriers (odds ratio [OR], 0.41; 95% confidence interval [CI], 0.20–0.84).

Although the frequency of the A allele at position -607 in inactive HBV carriers seems to be higher compared with those of CPLD patients (inactive HBV carrier; 64.0% vs CPLD; 54.0%), no significant difference was found between the inactive HBV carriers and the CPLD groups (Table VIII). However, the frequency of the C allele at position -137 was found to be significantly higher in the inactive HBV-carrier group compared with that in the CPLD group (Table VIII, inactive HBV carrier, 17.4% vs CPLD, 8.4%, $P = 0.024$).

To elucidate the relationship between these polymorphisms and the fibrosis staging, we divided the patients who received liver biopsy into 2 groups by the degree of fibrosis staging (the F0–F2 group, $n = 21$, and the F3–F4 group, $n = 36$). No significant difference was

Table VI. Genotype frequencies in patients with HBV

Genotype	Patients with HBV			
	Inactive HBV carrier (n = 43) (%)	CH (n = 62) (%)	Liver cirrhosis (n = 52) (%)	HCC (n = 47) (%)
Locus -607				
CC	6 (14.0)	6 (9.7)	10 (19.2)	8 (17.0)
CA	19 (44.2)	42 (67.7)	32 (61.5)	26 (55.3)
AA	18 (41.9)	14 (22.6)	10 (19.2)	13 (27.7)
Locus -137				
GG	31 (72.1)	53 (85.5)	40 (76.9)	43 (91.5)
GC	9 (20.9)	9 (14.5)	11 (21.2)	3 (6.4)
CC	3 (7.0)	0 (0)	1 (1.9)	1 (2.1)

Table VII. The distribution of IL-18 genotype in inactive HBV carriers and CPLD patients

Genotype	Inactive HBV Carrier (n = 43)		CPLD (n = 161)		OR (95% CI)	P value
	n	%	n	%		
Genotype -607						
A/A	18	41.9	37	23.0	0.41 (0.20–0.84)	0.022
A/C	19	44.2	100	62.1	2.07 (1.05–4.09)	0.051
C/C	6	14.0	24	14.9	1.08 (0.41–2.84)	>0.999
Genotype -137						
G/G	31	72.1	136	84.5	2.11 (0.95–4.65)	0.099
G/C	9	20.9	23	14.3	0.63 (0.27–1.48)	0.407
C/C	3	7.0	2	1.2	0.17 (0.03–1.04)	0.108

NOTES: P value: The Fisher exact test.

discovered in IL-18 gene-promoter polymorphisms (-607 or -137) between these 2 groups (Table IX).

DISCUSSION

The susceptibility to persistent HBV infection is governed by several factors, which include the age at infection. When infection is acquired during the early neonatal period from an HBV-infected mother, only 10% of children will eliminate the virus. In contrast, when infection is acquired during childhood or later, up to 90% will eliminate the virus spontaneously.¹⁶ Twin studies indicate that in addition to the age at infection, the host genetic background influences the outcome of HBV infection.³ Elimination of HBV infection requires an innate and adaptive humoral and cell-mediated immune response.¹⁷

Previous studies have shown that the capacity of cytokine production varies among individuals and correlates with the polymorphisms in the promoter region of